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☐ **1: GI "3650064" [GenBank]** Homo sapiens chromosome 17,... Related Sequences.

LOCUS AC004099 136222 bp DNA FFI 06-JUN-2000
 DEFINITION Homo sapiens chromosome 17, clone HCIT421K24, complete sequence.
 ACCESSION AC004099
 VERSION AC004099.1 GI:3650064
 KEYWORDS HTG
 SOURCE human.
 ORGANISM Homo sapiens
 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
 Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
 REFERENCE 1 (bases 1 to 136222)
 AUTHORS Birren,B., Linton,L., Nusbaum,C. and Lander,E.
 TITLE Homo sapiens chromosome 17, clone HCIT421K24
 JOURNAL Unpublished
 REFERENCE 2 (bases 1 to 136222)
 AUTHORS Birren,B., Fasman,K., McKernan,K., Nusbaum,C., Richardson,P.,
 Lander,E., Allen,N., Baker,J., Baldwin,J., Barna,N., Beckerly,R.,
 Boutwell,C., Byrne,S., Cantu,C., Castle,A., Cooke,P., Daly,M.J.,
 Depayre,E., Devon,K., Dewar,K., DuRette,B., Ettemadi,S.,
 Ferreira,P., Forrest,C., Gage,D., Gardyna,S., Gensheimer,S.,
 Geraigery,K., Gilmartin,T., Gray,D., Hedges,B., Harris,K.,
 Horton,L., Howland,J.C., Hui,L., Jostot,L., Linton,L.,
 MacKenzie,J., Marquis,N., McEwan,P., McGuirk,A., Meldrum,J.,
 Molla,M., Morris,W., Morrow,J., Nickman,A., Naylor,J., O'Connor,T.,
 Pavlin,B., Peterson,K., Ranganath,S., Riley,R., Roberts,D.,
 Rollins,G., Rossello,R., Roy,A., Shyam,R., Soohoo,S., Stilwell,J.,
 Stone,C., Strickland,C., Sydney,K., Tang,L., Vassiliev,H., Vo,A.,
 Warner,A., Wheeler,J., Wu,Y., Ye,W.J., Zemtseva,I., Zhao,J. and
 Zody,M.
 TITLE Direct Submission
 JOURNAL Submitted (01-FEB-1998) Whitehead Institute MIT Center for Genome
 Research, 20 Charles Street, Cambridge, MA 02141, USA
 REFERENCE 3 (bases 1 to 136222)
 AUTHORS Birren,B., Fasman,K., Linton,L., Nusbaum,C., Lander,E., Allen,N.,
 Baker,J., Baldwin,J., Barna,N., Beckerly,R., Benn,J., Boatin,C.,
 Boutwell,C., Brown,A., Byrne,S., Cantu,C., Castle,A., Cerny,J.,
 Cooke,P., Depayre,E., Devon,K., Dewar,K., Donelan,L., DuRette,B.,
 Ettemadi,S., Ferreira,P., FitzHugh,W., Forrest,C., Funke,B.,
 Gage,D., Geraigery,K., Gensheimer,S., Gilmartin,T.,
 Gray,D., Horton,L., Howland,J.C., Hui,L., Jostot,L., Linton,L.,
 MacKenzie,J., Marquis,N., McEwan,P., McGuirk,A., Meldrum,J.,
 Molla,M., Morris,W., Morrow,J., Nickman,A., Naylor,J., O'Connor,T.,
 Pavlin,B., Peterson,K., Ranganath,S., Riley,R., Roberts,D.,
 Rollins,G., Rossello,R., Roy,A., Shyam,R., Soohoo,S., Stilwell,J.,
 Stone,C., Strickland,C., Sydney,K., Tang,L., Vassiliev,H., Vo,A.,
 Warner,A., Wheeler,J., Wu,Y., Ye,W.J., Zemtseva,I., Zhao,J.

TITLE Warner, A., Wang, P., Wheeler, J., Wu, Y., Ye, W.J., Zhao, J. and Zody, M.
 JOURNAL Direct Submission
 REFERENCE Submitted (09-MAY-1998) Whitehead Institute/MIT Center for Genome
 AUTHORS Research, 320 Charles Street, Cambridge, MA 02141, USA
 4 (bases 1 to 136221)
 TITLE Warner, A., Wang, P., Wheeler, J., Wu, Y., Ye, W.J., Zhao, J. and Zody, M.
 JOURNAL Direct Submission
 REFERENCE Submitted (25-SEP-1998) Whitehead Institute/MIT Center for Genome
 AUTHORS Research, 320 Charles Street, Cambridge, MA 02141, USA
 5 (bases 1 to 136212)
 TITLE Warner, A., Wang, P., Wheeler, J., Wu, Y., Ye, W.J., Zhao, J. and Zody, M.
 JOURNAL Direct Submission
 REFERENCE Submitted (06-JUN-2000) Whitehead Institute/MIT Center for Genome
 COMMENT Research, 320 Charles Street, Cambridge, MA 02141, USA
 On Sep 25, 1998 this sequence version replaced gi:3126780.
 All repeats were identified using RepeatMasker: Smit, A.F.A. &
 Green, P. (1996-1997)
<http://ftp.genome.washington.edu/EM/RepeatMasker.html>.
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 <map="17"
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repeat_region complement (1626..1667)
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repeat_region 3619..3673
/rpt_family="Alu"
repeat_region complement (3692..4000)
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repeat_region 12531..12814
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repeat_region 12816..13111
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repeat_region      complement(20815..21109)
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repeat_region      complement(21164..21338)
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repeat_region      complement(22398..22699)
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repeat_region      complement(22751..22958)
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repeat_region      23101..23425
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                    /note="Single-stranded coverage."
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feature            59944..59950
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                    /rpt_family="MIE"
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repeat_region      complement(101611..101782)
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repeat_region      complement(101786..102086)
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repeat_region complement(102309..102607)
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repeat_region 102608..102717
/rpt_family "MER63"
repeat_region 104414..104441
/rpt_family "AT-rich"
repeat_region 104560..104636
/rpt_family "MLT11"
repeat_region 104697..104834
/rpt_family "AluSx"
repeat_region 104835..105128
/rpt_family "AluSq"
repeat_region 105129..105293
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repeat_region 105294..105550
/rpt_family "MLT11"
repeat_region complement(105628..106079)
/rpt_family "MLT10"
repeat_region complement(106661..106987)
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repeat_region 107204..107329
/rpt_family "MIR"
repeat_region complement(107344..107524)
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/rpt_family "(TA)n"
repeat_region 108471..108690
/rpt_family "L1ME1"
repeat_region 109398..109473
/rpt_family "MIR"
repeat_region 110116..110383
/rpt_family "AluJb"
repeat_region 110641..110944
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Figure 1. The effect of the number of trials on the number of correct responses. The number of correct responses was significantly higher than the number of incorrect responses in all cases. The number of correct responses was significantly higher than the number of incorrect responses in all cases. The number of correct responses was significantly higher than the number of incorrect responses in all cases.

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Figure 1. The effect of the number of trials on the number of correct responses. The number of correct responses was significantly higher than the number of incorrect responses in all cases. The number of correct responses was significantly higher than the number of incorrect responses in all cases. The number of correct responses was significantly higher than the number of incorrect responses in all cases.

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